Table 2. Pairs of primer sequences used for SSCP analysis of *BLM*.

Name	Forward sequence[a]	SEQ. ID. NO.	Reverse sequence[a]	SEQ. ID. NO.	Product length (bp)
Cl-B	GGATCCTCGGTTCCGTCCGC	(10)	GAGGTTCACTGAAGGAAAAGTTC	(26)	269
Cl-A	CAACTAGAACGTCACTCAGCC	(11)	GAAGTCCTTGACCCTTGCTG	(27)	233
Cl-1	GACTTTTCCTTCAGTGAACCTC	(12)	GGGATTTCTTTACAGTTGGTGTG	(28)	186
Cl-2	CCAGATTTCTTGCAGACTCCG	(13)	CTCTTACAAAGTGAACTTGGGGG	(29)	213
Cl-3	CTTTAAGTACCATCAATGATTGGG	(14)	CCTCAGTCAAATCTATTTGCTCG	(30)	227
Cl-4	GAGTAAGCACTGCTCAGAAATC	(15)	GCTTAACCATTCTGAGTCATCC	(31)	160
Cl-5	CGAGCAAATAGATTTGACTGAGG	(16)	CAATACATGGAACTTTCTCAGTTG	(32)	223
Cl-6	GAAGATCTCAGGAAAGTGAC	(17)	CGTACTAAGGCATTTTGAAGAGG	(33)	215
Cl-7	CAACTGAGAAAGTTCCATGTATTG	(18)	CACAGTCTGTGCTGGTTTCTG	(34)	239
Cl-9	CTATTCCTGATGATAAACTGAAAC	(19)	CCTTCATAGAATTCCCTGTAGG	(35)	200
Cl-10	GTGGGAGATACAGGCCTGATTC	(20)	GTGTTTCAGCCCAGTTGCTAC	(36)	244
Cl-11	CAGGATTTCTCTGCCACCAGG	(21)	GCAGTATGTTTATTCTGATCTTTC	(37)	183
Cl-12	CAGGAAATGTTTCTCACAAGCAC	(22)	CCTTGATGGGGTTGATAGGCAG	(38)	203
Cl-13	CAGCCAGCAAATCTTCCACAG	(23)	CGCTTCCATGTTTCAGATTTCTGG	(39)	204
Cl-14	GAATTATACTGACAAGTCAGCAC	(24)	GATCTACGATAAGTGATCTCAAG	(40)	295
Cl-15	CTCCTGGGGTCACTGTTGTC	(25)	GAGTCTGTTACTTGCACAGATC	(41)	211
Cl-16	CAATCATAAAACTTCTATATGTCAC	(42)	GCCATCACCGGAACACAGAAGG	(57)	207
Cl-17	GTGGGGACATGATTTTCGTCAAG	(43)	GATTATGTCTGTTAAAGCTCATG	(58)	175
Cl-18	GACATCCTGACTCAGCTGAAG	(44)	CGTGTCAGCCATGGTGTCAC	(59)	203
Cl-19	GCACCACCCATATGAATTCAGG	(45)	CAGATAACCTGACAGCCATCC	(60)	179
Cl-20	GATGAAGTGCAGCAGAAGTGG	(46)	CAGTGTCGGTCACATCATGATAG	(61)	221
Cl-21	GCAGAGACTGGAAGAGATGGG	(47)	GCTGTATTCTCCTGCATTCCG	(62)	188
Cl-22	GTATAGCATGGTACATTACTGTG	(48)	CCTGTGATGAACTATGTTCTTG	(63)	228
Cl-23	GACTGACGATGTGAAAAGTATTG	(49)	CCAAAATCTGTCAAGTATCAGC	(64)	235
Cl-24	CCAGTCAGGTATATTTGGAAAAG	(50)	GGAATTTTCTGTTTCCATAAAGTC	(65)	206
Cl-25	CGATCGCCTTATGTGATGCTCG	(51)	CAAGCTTCTTGAGAGTGACGG	(66)	248
Cl-26	GAACTTACAGAAGTCTGCAAATC	(52)	GATGTCCATTCCTCGTCAAGCTC	(67)	208
Cl-27	GGTGTTACTGAAGAACAAACTGG	(53)	GGGTATTCCTCGTCAAGCTC	(68)	168
Cl-28	GGATAAGCCTGTCCAGCAGC	(54)	CCTAGATATCTTTCTACATGTGG	(69)	214
Cl-29	GCTTCCAGTGGTTCCAAGGC	(55)	GTTATGAGAATGCATATGAAGGC	(70)	204
Cl-30	CTCAAGCGCGACATCAGGAGCC	(56)	CAAGAATAACAGCTTTATAGTCAC	(71)	178

[a] 5' to 3'